

Bis(2-amino-3-carboxypyrazin-1-i um)sulfate dihydrate

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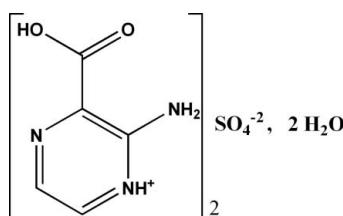
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Key indicators: single-crystal X-ray study; $T = 150\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$; R factor = 0.035; wR factor = 0.096; data-to-parameter ratio = 14.9.

The crystal structure of the title compound, $2\text{C}_5\text{H}_6\text{N}_3\text{O}_2^+ \cdot \text{SO}_4^{2-} \cdot 2\text{H}_2\text{O}$, displays a variety of $\text{N}-\text{H}\cdots\text{O}$ and $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds in which all potential donors and acceptors are involved. In the crystal, cations and anions are interconnected, forming $R_3^3(10)$ and $R_2^2(8)$ ring motifs whereas the anions and water molecules form $R_2^2(10)$ rings, which develop in chains running along [100]. The resulting three-dimensional network exhibits undulating sheets parallel to (011), marked by the presence of $R_6^6(26)$ rings in which six cations are involved.

Related literature

For related compounds, see: Berrah *et al.* (2005a,b, 2011); Bouacida *et al.* (2005, 2009); Dobson & Gerkin (1996). For hydrogen-bond motifs, see: Bernstein *et al.* (1995); Etter *et al.* (1990). For similar intermolecular interactions, see: Dorn *et al.* (2005), Janiak (2000); Desiraju (2003).



Experimental

Crystal data

$2\text{C}_5\text{H}_6\text{N}_3\text{O}_2^+ \cdot \text{SO}_4^{2-} \cdot 2\text{H}_2\text{O}$	$V = 1605.36 (17)\text{ \AA}^3$
$M_r = 412.36$	$Z = 4$
Monoclinic, $P2_1/a$	Mo $K\alpha$ radiation
$a = 7.7214 (4)\text{ \AA}$	$\mu = 0.27\text{ mm}^{-1}$
$b = 20.7043 (14)\text{ \AA}$	$T = 150\text{ K}$
$c = 10.6398 (7)\text{ \AA}$	$0.55 \times 0.36 \times 0.15\text{ mm}$
$\beta = 109.299 (2)^{\circ}$	

Data collection

Bruker APEXII diffractometer	13466 measured reflections
Absorption correction: multi-scan (<i>SADABS</i> ; Sheldrick, 2002)	3675 independent reflections
$(SADABS$; Sheldrick, 2002)	3146 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.708$, $T_{\max} = 0.960$	$R_{\text{int}} = 0.039$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.035$	246 parameters
$wR(F^2) = 0.096$	H-atom parameters constrained
$S = 1.03$	$\Delta\rho_{\max} = 0.47\text{ e \AA}^{-3}$
3675 reflections	$\Delta\rho_{\min} = -0.48\text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (\AA , $^{\circ}$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N1A—H1A1…O4	0.88	1.92	2.7970 (18)	175
N1A—H1A2…O6A	0.88	2.04	2.6741 (18)	128
N1A—H1A2…O6B ⁱ	0.88	2.30	3.0158 (18)	138
N2A—H2A…O1	0.88	1.83	2.6915 (18)	167
N1B—H1B1…O2	0.88	2.34	3.0827 (18)	142
N1B—H1B2…O6B	0.88	2.08	2.7144 (19)	129
N1B—H1B2…O6A ⁱⁱ	0.88	2.10	2.8237 (18)	139
N2B—H2B…O1W	0.88	1.81	2.6705 (18)	167
O5B—H5B…O2W	0.84	1.67	2.5046 (17)	174
O5A—H5A…O2 ⁱ	0.84	1.78	2.6192 (16)	175
O1W—H1W…O3	0.85	1.95	2.7934 (19)	175
O1W—H2W…O2 ⁱⁱⁱ	0.85	2.06	2.8996 (18)	167
O1W—H2W…O4 ⁱⁱⁱ	0.85	2.65	3.2856 (17)	133
O2W—H3W…O4 ⁱⁱ	0.85	1.88	2.7351 (17)	177
O2W—H4W…O3 ^{iv}	0.86	1.91	2.7633 (17)	171

Symmetry codes: (i) $x, y, z - 1$; (ii) $x, y, z + 1$; (iii) $x + \frac{1}{2}, -y + \frac{1}{2}, z$; (iv) $x - \frac{1}{2}, -y + \frac{1}{2}, z + 1$.

Data collection: *APEx2* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SIR2002* (Burla *et al.*, 2003); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997) and *DIAMOND* (Bränenburg & Berndt, 2001); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: DN2656).

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supplementary materials

Acta Cryst. (2011). E67, o677-o678 [doi:10.1107/S1600536811005824]

Bis(2-amino-3-carboxypyrazin-1-i um) sulfate dihydrate

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Comment

Hydrogen bonds are object of several studies, that aim at elucidate their influence on crystal construction and compounds properties (Desiraju, 2003). N-heterocyclic compounds such as pyrazine and its derivatives may be interesting units to built new edifices involving original hydrogen-bonding scheme since they include a variety of potential hydrogen donors and acceptors. In this perspective and as a part of our search for new hybrid compounds based on protonated amines and imines (Berrah *et al.* 2011, 2005*a,b*; Bouacida *et al.* 2005,2009), we present here the structure of Bis (2-Amino-3-carboxypyrazin-1-i um) sulfate dihydrate.

The asymmetric units of (I) includes two symmetry- independent cations and water molecules, and one sulfate anion. Cations and anions are interconnected to form $R_3^3(10)$ and $R_2^2(8)$ ring motifs (Etter *et al.*, 1990; Bernstein *et al.*, 1995) (Fig 1). Bond lengths and angles are as expected (Berrah *et al.* 2011; Dobson & Gerkin, 1996).

The three-dimensional structure of (I), results from undulating sheets of cations dimmers parallel to (011) plane(Fig.2 and Fig.3)and sulfate-water chains extending along [100](Fig.3). An interesting hydrogen bonds network, in which all potential donors and acceptors are involved, and especially marked by the presence of $R_6^6(26)$ and $R_2^3(10)$ set-graph motifs (Etter *et al.*, 1990; Bernstein *et al.*, 1995), ensures the coherence of the structure(Fig.2 and Fig.3, table 1). This later is reinforced by the contribution of $\pi\cdots\pi$, S—O $\cdots\pi$ and C—O $\cdots\pi$ interactions (Dorn *et al.* 2005; Janiak, 2000) (table 2,3).

Experimental

The title compound was synthesized by reacting 3-amino-pyrazine 2- carboxylic acid with some excess of sulphuric acid in aqueous solution. Slow evaporation leads to well crystallized yellow needles.

Refinement

All non-H atoms were refined with anisotropic atomic displacement parameters. H atoms of water molecule were located in difference Fourier maps and treated as riding on their parent oxygen atoms with O—H = 0.85, H \cdots H = 1.40 and $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$. The remaining H atoms were localized on Fourier maps but introduced in calculated positions and treated as riding on their parent atoms (C, N or O) with C—H = 0.95 Å, O—H = 0.84 Å and N—H = 0.88 Å with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}$ or N) and $U_{\text{iso}}(\text{H} = 1.5 U_{\text{eq}}(\text{O})$.

supplementary materials

Figures

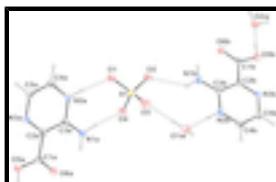


Fig. 1. The asymmetric unit of the title compound with the atomic labelling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are represented as small spheres of arbitrary radii. Hydrogen bonds are shown as dashed lines.

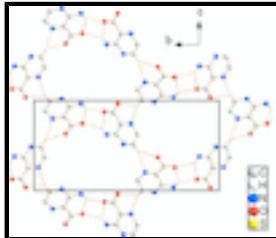


Fig. 2. Partial packing view showing undulating sheets parallel to (011) plane and $R_6^6(26)$ rings set motif. Hydrogen bonds are shown as dashed lines. Hydrogen atoms not involved in H-bonds have been omitted for clarity

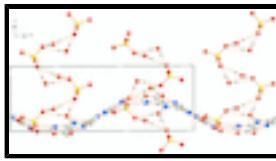


Fig. 3. Partial packing view showing sulfate-water chains extending along [100] direction and undulating sheets. Hydrogen bonds are shown as dashed lines. Hydrogen atoms not involved in H-bonds have been omitted for clarity.

Bis(2-amino-3-carboxypyrazin-1-ium) sulfate dihydrate

Crystal data

$2\text{C}_5\text{H}_6\text{N}_3\text{O}_2^+\cdot\text{SO}_4^{2-}\cdot2\text{H}_2\text{O}$	$F(000) = 856$
$M_r = 412.36$	$D_x = 1.706 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/a$	Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
$a = 7.7214 (4) \text{ \AA}$	Cell parameters from 5179 reflections
$b = 20.7043 (14) \text{ \AA}$	$\theta = 2.8\text{--}27.5^\circ$
$c = 10.6398 (7) \text{ \AA}$	$\mu = 0.27 \text{ mm}^{-1}$
$\beta = 109.299 (2)^\circ$	$T = 150 \text{ K}$
$V = 1605.36 (17) \text{ \AA}^3$	Prism, yellow
$Z = 4$	$0.55 \times 0.36 \times 0.15 \text{ mm}$

Data collection

Bruker APEXII diffractometer	3146 reflections with $I > 2\sigma(I)$
graphite	$R_{\text{int}} = 0.039$
CCD rotation images, thin slices scans	$\theta_{\text{max}} = 27.6^\circ, \theta_{\text{min}} = 2.3^\circ$
Absorption correction: multi-scan (<i>SADABS</i> ; Sheldrick, 2002)	$h = -8 \rightarrow 9$
$T_{\text{min}} = 0.708, T_{\text{max}} = 0.960$	$k = -26 \rightarrow 26$
13466 measured reflections	$l = -13 \rightarrow 13$
3675 independent reflections	

Refinement

Refinement on F^2	Primary atom site location: structure-invariant direct methods
Least-squares matrix: full	Secondary atom site location: difference Fourier map
$R[F^2 > 2\sigma(F^2)] = 0.035$	Hydrogen site location: inferred from neighbouring sites
$wR(F^2) = 0.096$	H-atom parameters constrained
$S = 1.03$	$w = 1/[\sigma^2(F_o^2) + (0.0491P)^2 + 0.7394P]$ where $P = (F_o^2 + 2F_c^2)/3$
3675 reflections	$(\Delta/\sigma)_{\max} = 0.001$
246 parameters	$\Delta\rho_{\max} = 0.47 \text{ e } \text{\AA}^{-3}$
0 restraints	$\Delta\rho_{\min} = -0.48 \text{ e } \text{\AA}^{-3}$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) etc. and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
O5A	0.17659 (16)	0.04617 (6)	0.03501 (11)	0.0221 (3)
H5A	0.1515	0.0732	-0.0273	0.033*
O6A	0.03846 (18)	0.11842 (6)	0.12557 (11)	0.0258 (3)
N1A	0.0567 (2)	0.10587 (7)	0.37979 (13)	0.0231 (3)
H1A1	0.0399	0.1167	0.4549	0.028*
H1A2	0.0122	0.1303	0.3087	0.028*
N2A	0.21592 (19)	0.01604 (6)	0.48490 (13)	0.0195 (3)
H2A	0.1989	0.0288	0.5588	0.023*
N3A	0.27826 (19)	-0.02410 (6)	0.25952 (13)	0.0194 (3)
C1A	0.1266 (2)	0.06905 (7)	0.13275 (15)	0.0185 (3)
C2A	0.1875 (2)	0.02980 (7)	0.25829 (15)	0.0177 (3)
C3A	0.1485 (2)	0.05305 (7)	0.37379 (15)	0.0183 (3)
C4A	0.3079 (2)	-0.03946 (8)	0.48622 (16)	0.0207 (3)
H4A	0.3518	-0.0644	0.5654	0.025*
C5A	0.3374 (2)	-0.05941 (8)	0.37264 (16)	0.0214 (3)
H5C	0.4006	-0.0988	0.3731	0.026*
O5B	-0.03794 (19)	0.32977 (6)	1.37046 (12)	0.0273 (3)
H5B	-0.0768	0.3055	1.4185	0.041*

supplementary materials

O6B	-0.03786 (17)	0.23824 (5)	1.25812 (11)	0.0241 (3)
N1B	0.0619 (2)	0.24559 (7)	1.03668 (14)	0.0238 (3)
H1B1	0.0821	0.2313	0.9649	0.029*
H1B2	0.0315	0.2184	1.0895	0.029*
N2B	0.12393 (19)	0.34886 (7)	0.98346 (13)	0.0205 (3)
H2B	0.1422	0.3332	0.912	0.025*
N3B	0.07073 (19)	0.39911 (6)	1.20349 (13)	0.0206 (3)
C1B	-0.0126 (2)	0.29617 (8)	1.27448 (15)	0.0196 (3)
C2B	0.0482 (2)	0.33663 (7)	1.17934 (14)	0.0178 (3)
C3B	0.0773 (2)	0.30786 (8)	1.06522 (15)	0.0187 (3)
C4B	0.1438 (2)	0.41269 (8)	1.00633 (16)	0.0227 (3)
H4B	0.1752	0.4403	0.9458	0.027*
C5B	0.1180 (2)	0.43737 (8)	1.11810 (16)	0.0233 (3)
H5D	0.1338	0.4824	1.1357	0.028*
S1	0.15317 (5)	0.131125 (18)	0.73117 (4)	0.01813 (11)
O1	0.2146 (2)	0.06509 (6)	0.71883 (12)	0.0337 (3)
O2	0.09536 (18)	0.13565 (6)	0.85065 (12)	0.0262 (3)
O3	0.30149 (17)	0.17773 (7)	0.74478 (12)	0.0306 (3)
O4	-0.00482 (16)	0.14708 (6)	0.61180 (11)	0.0236 (3)
O1W	0.23309 (17)	0.30737 (6)	0.78406 (12)	0.0262 (3)
H1W	0.2465	0.2676	0.7697	0.039*
H2W	0.3352	0.3253	0.7903	0.039*
O2W	-0.15818 (19)	0.26453 (6)	1.52193 (12)	0.0296 (3)
H3W	-0.1068	0.2285	1.5511	0.044*
H4W	-0.1653	0.2863	1.5886	0.044*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O5A	0.0332 (7)	0.0213 (6)	0.0140 (5)	0.0044 (5)	0.0109 (5)	0.0030 (4)
O6A	0.0400 (7)	0.0206 (6)	0.0184 (6)	0.0076 (5)	0.0115 (5)	0.0036 (4)
N1A	0.0340 (8)	0.0227 (7)	0.0140 (6)	0.0063 (6)	0.0100 (6)	0.0012 (5)
N2A	0.0269 (7)	0.0193 (6)	0.0128 (6)	-0.0002 (5)	0.0072 (5)	0.0001 (5)
N3A	0.0253 (7)	0.0168 (6)	0.0158 (6)	-0.0013 (5)	0.0063 (5)	-0.0003 (5)
C1A	0.0232 (8)	0.0172 (7)	0.0153 (7)	-0.0026 (6)	0.0065 (6)	-0.0007 (6)
C2A	0.0225 (8)	0.0159 (7)	0.0154 (7)	-0.0020 (6)	0.0072 (6)	-0.0008 (6)
C3A	0.0229 (8)	0.0175 (7)	0.0144 (7)	-0.0027 (6)	0.0062 (6)	-0.0007 (6)
C4A	0.0259 (8)	0.0177 (7)	0.0168 (7)	-0.0016 (6)	0.0048 (6)	0.0035 (6)
C5A	0.0279 (9)	0.0163 (7)	0.0186 (7)	0.0011 (6)	0.0056 (6)	0.0000 (6)
O5B	0.0480 (8)	0.0198 (6)	0.0201 (6)	-0.0009 (5)	0.0194 (5)	-0.0004 (5)
O6B	0.0355 (7)	0.0185 (6)	0.0193 (6)	-0.0015 (5)	0.0104 (5)	0.0003 (4)
N1B	0.0372 (8)	0.0188 (7)	0.0167 (6)	-0.0035 (6)	0.0105 (6)	-0.0037 (5)
N2B	0.0267 (7)	0.0221 (7)	0.0128 (6)	-0.0027 (5)	0.0069 (5)	-0.0024 (5)
N3B	0.0267 (7)	0.0172 (6)	0.0167 (6)	0.0005 (5)	0.0056 (5)	-0.0008 (5)
C1B	0.0231 (8)	0.0198 (8)	0.0141 (7)	0.0019 (6)	0.0039 (6)	0.0008 (6)
C2B	0.0216 (8)	0.0173 (7)	0.0127 (7)	0.0005 (6)	0.0033 (6)	0.0001 (5)
C3B	0.0203 (8)	0.0194 (7)	0.0142 (7)	-0.0010 (6)	0.0029 (6)	-0.0014 (6)
C4B	0.0276 (9)	0.0201 (8)	0.0195 (7)	-0.0035 (6)	0.0068 (6)	0.0020 (6)

C5B	0.0324 (9)	0.0177 (7)	0.0193 (7)	-0.0019 (6)	0.0077 (7)	0.0004 (6)
S1	0.0259 (2)	0.01758 (19)	0.01276 (18)	0.00290 (14)	0.00883 (15)	0.00224 (13)
O1	0.0605 (9)	0.0245 (6)	0.0186 (6)	0.0188 (6)	0.0163 (6)	0.0047 (5)
O2	0.0429 (7)	0.0230 (6)	0.0193 (6)	0.0051 (5)	0.0192 (5)	0.0035 (5)
O3	0.0291 (7)	0.0384 (7)	0.0238 (6)	-0.0067 (5)	0.0080 (5)	0.0029 (5)
O4	0.0266 (6)	0.0246 (6)	0.0179 (6)	0.0031 (5)	0.0049 (5)	0.0017 (4)
O1W	0.0294 (6)	0.0273 (6)	0.0252 (6)	-0.0009 (5)	0.0136 (5)	-0.0005 (5)
O2W	0.0460 (8)	0.0244 (6)	0.0229 (6)	0.0086 (5)	0.0174 (6)	0.0061 (5)

Geometric parameters (\AA , $^\circ$)

O5A—C1A	1.3116 (19)	N1B—H1B1	0.88
O5A—H5A	0.84	N1B—H1B2	0.88
O6A—C1A	1.216 (2)	N2B—C4B	1.343 (2)
N1A—C3A	1.316 (2)	N2B—C3B	1.347 (2)
N1A—H1A1	0.88	N2B—H2B	0.88
N1A—H1A2	0.88	N3B—C2B	1.319 (2)
N2A—C4A	1.349 (2)	N3B—C5B	1.344 (2)
N2A—C3A	1.360 (2)	C1B—C2B	1.503 (2)
N2A—H2A	0.88	C2B—C3B	1.435 (2)
N3A—C2A	1.315 (2)	C4B—C5B	1.368 (2)
N3A—C5A	1.352 (2)	C4B—H4B	0.95
C1A—C2A	1.500 (2)	C5B—H5D	0.95
C2A—C3A	1.442 (2)	S1—O1	1.4670 (12)
C4A—C5A	1.365 (2)	S1—O3	1.4677 (13)
C4A—H4A	0.95	S1—O4	1.4786 (12)
C5A—H5C	0.95	S1—O2	1.4831 (12)
O5B—C1B	1.3028 (19)	O1W—H1W	0.8491
O5B—H5B	0.84	O1W—H2W	0.8542
O6B—C1B	1.218 (2)	O2W—H3W	0.8543
N1B—C3B	1.321 (2)	O2W—H4W	0.8582
C1A—O5A—H5A	109.5	C4B—N2B—C3B	122.81 (14)
C3A—N1A—H1A1	120	C4B—N2B—H2B	118.6
C3A—N1A—H1A2	120	C3B—N2B—H2B	118.6
H1A1—N1A—H1A2	120	C2B—N3B—C5B	119.63 (14)
C4A—N2A—C3A	122.57 (14)	O6B—C1B—O5B	125.38 (15)
C4A—N2A—H2A	118.7	O6B—C1B—C2B	121.55 (14)
C3A—N2A—H2A	118.7	O5B—C1B—C2B	113.05 (14)
C2A—N3A—C5A	119.32 (14)	N3B—C2B—C3B	121.62 (14)
O6A—C1A—O5A	123.99 (14)	N3B—C2B—C1B	117.76 (14)
O6A—C1A—C2A	121.03 (14)	C3B—C2B—C1B	120.62 (14)
O5A—C1A—C2A	114.98 (13)	N1B—C3B—N2B	119.31 (15)
N3A—C2A—C3A	122.39 (14)	N1B—C3B—C2B	124.91 (15)
N3A—C2A—C1A	118.54 (14)	N2B—C3B—C2B	115.78 (14)
C3A—C2A—C1A	119.06 (14)	N2B—C4B—C5B	118.98 (15)
N1A—C3A—N2A	118.94 (14)	N2B—C4B—H4B	120.5
N1A—C3A—C2A	125.91 (14)	C5B—C4B—H4B	120.5
N2A—C3A—C2A	115.15 (14)	N3B—C5B—C4B	121.17 (15)
N2A—C4A—C5A	119.33 (14)	N3B—C5B—H5D	119.4

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N2A—C4A—H4A	120.3	C4B—C5B—H5D	119.4
C5A—C4A—H4A	120.3	O1—S1—O3	110.91 (9)
N3A—C5A—C4A	121.21 (15)	O1—S1—O4	109.31 (7)
N3A—C5A—H5C	119.4	O3—S1—O4	109.46 (7)
C4A—C5A—H5C	119.4	O1—S1—O2	109.47 (7)
C1B—O5B—H5B	109.5	O3—S1—O2	108.66 (7)
C3B—N1B—H1B1	120	O4—S1—O2	109.00 (7)
C3B—N1B—H1B2	120	H1W—O1W—H2W	105.7
H1B1—N1B—H1B2	120	H3W—O2W—H4W	108
C5A—N3A—C2A—C3A	0.1 (2)	C5B—N3B—C2B—C3B	1.6 (2)
C5A—N3A—C2A—C1A	178.70 (14)	C5B—N3B—C2B—C1B	-177.43 (14)
O6A—C1A—C2A—N3A	178.18 (15)	O6B—C1B—C2B—N3B	179.35 (15)
O5A—C1A—C2A—N3A	-2.2 (2)	O5B—C1B—C2B—N3B	0.9 (2)
O6A—C1A—C2A—C3A	-3.2 (2)	O6B—C1B—C2B—C3B	0.3 (2)
O5A—C1A—C2A—C3A	176.41 (14)	O5B—C1B—C2B—C3B	-178.12 (14)
C4A—N2A—C3A—N1A	178.56 (15)	C4B—N2B—C3B—N1B	-179.42 (15)
C4A—N2A—C3A—C2A	-1.9 (2)	C4B—N2B—C3B—C2B	0.5 (2)
N3A—C2A—C3A—N1A	-179.04 (16)	N3B—C2B—C3B—N1B	178.17 (15)
C1A—C2A—C3A—N1A	2.4 (2)	C1B—C2B—C3B—N1B	-2.9 (2)
N3A—C2A—C3A—N2A	1.5 (2)	N3B—C2B—C3B—N2B	-1.7 (2)
C1A—C2A—C3A—N2A	-177.09 (13)	C1B—C2B—C3B—N2B	177.27 (14)
C3A—N2A—C4A—C5A	0.8 (2)	C3B—N2B—C4B—C5B	0.8 (2)
C2A—N3A—C5A—C4A	-1.4 (2)	C2B—N3B—C5B—C4B	-0.2 (3)
N2A—C4A—C5A—N3A	0.9 (2)	N2B—C4B—C5B—N3B	-1.0 (3)

Hydrogen-bond geometry (\AA , $^\circ$)

$D\cdots H$	$D—H$	$H\cdots A$	$D\cdots A$	$D—H\cdots A$
N1A—H1A1···O4	0.88	1.92	2.7970 (18)	175.
N1A—H1A2···O6A	0.88	2.04	2.6741 (18)	128.
N1A—H1A2···O6B ⁱ	0.88	2.30	3.0158 (18)	138.
N2A—H2A···O1	0.88	1.83	2.6915 (18)	167.
N1B—H1B1···O2	0.88	2.34	3.0827 (18)	142.
N1B—H1B2···O6B	0.88	2.08	2.7144 (19)	129.
N1B—H1B2···O6A ⁱⁱ	0.88	2.10	2.8237 (18)	139.
N2B—H2B···O1W	0.88	1.81	2.6705 (18)	167.
O5B—H5B···O2W	0.84	1.67	2.5046 (17)	174.
O5A—H5A···O2 ⁱ	0.84	1.78	2.6192 (16)	175.
O1W—H1W···O3	0.85	1.95	2.7934 (19)	175.
O1W—H2W···O2 ⁱⁱⁱ	0.85	2.06	2.8996 (18)	167
O1W—H2W···O4 ⁱⁱⁱ	0.85	2.65	3.2856 (17)	133.
O2W—H3W···O4 ⁱⁱ	0.85	1.88	2.7351 (17)	177.
O2W—H4W···O3 ^{iv}	0.86	1.91	2.7633 (17)	171.
C4A—H4A···O5B ^v	0.95	2.58	3.320 (2)	134.
C4A—H4A···N3B ^v	0.95	2.45	3.369 (2)	163.
C4B—H4B···O5A ^{vi}	0.95	2.45	3.187 (2)	134.
C4B—H4B···N3A ^{vi}	0.95	2.44	3.347 (2)	159

C5A—H5C···O1W ^{vii}	0.95	2.55	3.175 (2)	124.
C5B—H5D···O1 ^{viii}	0.95	2.35	3.192 (2)	148.
Symmetry codes: (i) $x, y, z-1$; (ii) $x, y, z+1$; (iii) $x+1/2, -y+1/2, z$; (iv) $x-1/2, -y+1/2, z+1$; (v) $-x+1/2, y-1/2, -z+2$; (vi) $-x+1/2, y+1/2, -z+1$; (vii) $-x+1/2, y-1/2, -z+1$; (viii) $-x+1/2, y+1/2, -z+2$.				

Table 2

π — π stacking interactions (\AA , °)

Cg1 is the centroid of the N2A—C4A ring.

CgI	CgJ	CgI···CgJ ^a	α	β	γ	CgI···P(J) ^b	CgJ···P(I) ^c	Slippage
Cg1	Cg1 ⁱ	3.9678 (9)	0	34.94	34.94	3.2528 (6)	3.2527 (6)	2.272

Symmetry codes: (i) $l-x, -y, l-z$. Notes: a : Distance between centroids b : Perpendicular distance of CgI on ring plan J c : Perpendicular distance of CgJ on ring plan I α = Dihedral Angle between the ring planes β = Angle between the centroid vector CgI···CgJ and the normal to the plane I. γ = Angle between the centroid vector CgI···CgJ and the normal to the plane J. Slippage = vertical displacement between ring centroids.

Table 3

S—O··· π and C—O··· π interactions (\AA , °).

Cg1 and Cg2 are the centroids of the N2A—C4A and N2B—C4B rings, respectively.

X	I	J	I···J	X—I···J	X···J
S1	O1	Cg1 ⁱ	3.5922 (17)	91.83 (7)	3.9233 (8)
S1	O2	Cg1 ⁱ	3.9845 (14)	76.88 (5)	3.9233 (8)
S1	O2	Cg2 ⁱⁱ	3.8831 (15)	92.77 (6)	4.2231 (8)
C1A	O6A	Cg2 ⁱⁱⁱ	3.3136 (16)	125.62 (11)	4.1418 (18)

Symmetry codes: (i) $-x, -y, 1-z$; (ii) $x, y, 1+z$; (iii) $x-1, y, z$.

supplementary materials

Fig. 1

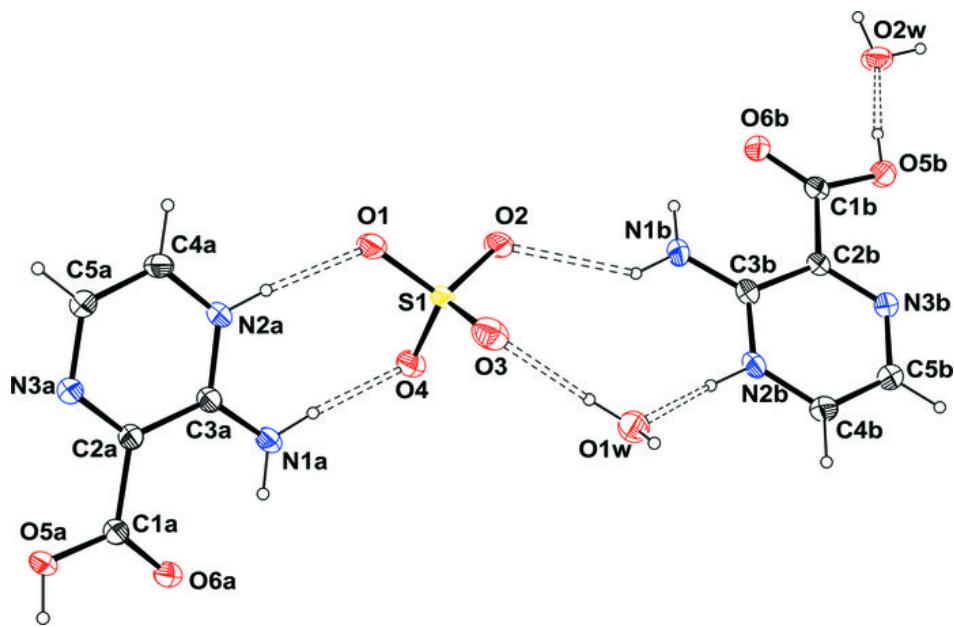
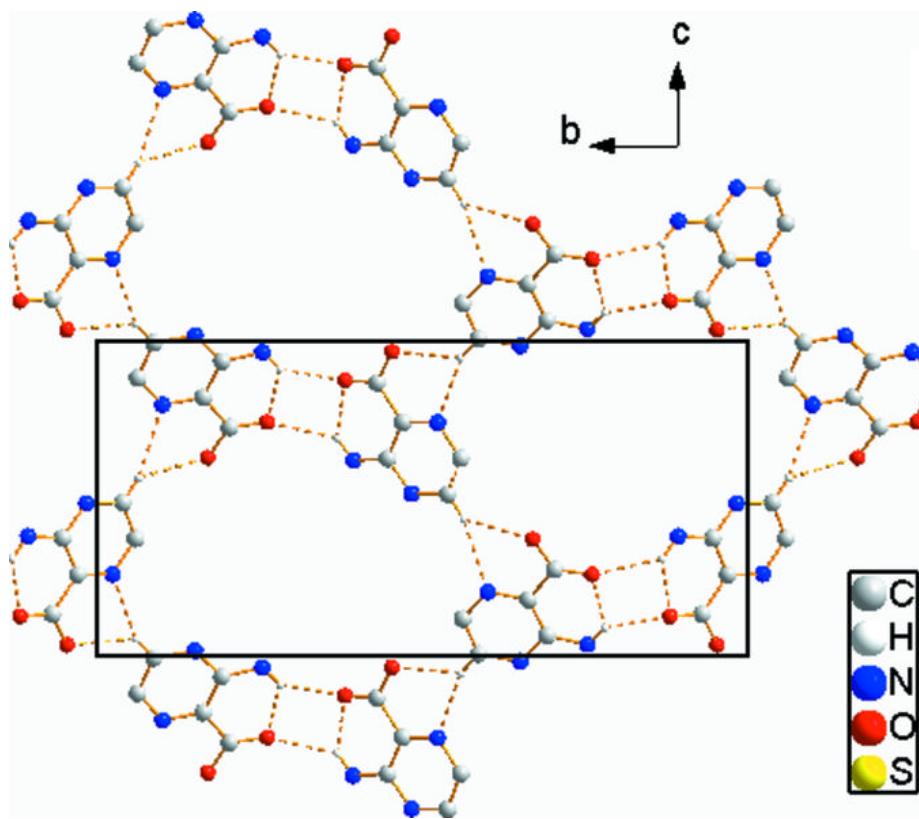


Fig. 2



C
H
N
O
S
P

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Fig. 3

